

XX  
 PD 24-JUN-2004.  
 XX  
 PP 05-DEC-2003; 2003WO-GB05323.  
 XX  
 PR 06-DEC-2002; 2002US-0431620P.  
 XX  
 PA (SIGE-) SINGAPORE GEN HOSPITAL PTE LTD.  
 PA (DENI/) DENISON C M.  
 XX  
 PT Xiao Z;  
 XX DR WPI; 2004-468811/44.  
 DR p-PSDB; ADQ16419.  
 XX  
 PT New peptides that interact with myelin proteins Nogo, TNR and MAG, useful in preparing a composition for treating CNS damage, spinal cord injury or stroke.  
 XX PS Disclosure; SEQ ID NO 16; 81pp; English.  
 CC The present sequence encodes Nogo-66 domain b. The specification describes peptides which interact with the myelin proteins Nogo (specifically the Nogo-66 domain), the extracellular matrix glycoprotein tenascin-R (Tn-R) (specifically Tn-R epidermal growth factor like (TNR-EGR)) and myelin-associated glycoprotein (MAG). These proteins have neural growth inhibitory activity. The peptide is isolated from a 7-mer peptide display library exposed to a plate coated with the target protein. Peptides of the invention are useful in preparing a composition for treating central nervous system (CNS) damage, spinal cord injury or stroke. The peptides may also be used in vaccines against myelin antigens. The vaccine is based on the specific inhibitory portions of major myelin proteins, instead of the whole protein.  
 CC Sequence 198 BP; 56 A; 36 C; 49 G; 57 T; 0 U; 0 Other;  
 CC Query Match 100.0%; Score 25; DB 12; Length 198;  
 CC Best Local Similarity 72.0%; Pred. No. 0.079; Mismatches 0; Indels 0; Gaps 0;  
 CC Matches 18; Conservative 7; Mismatches 0; Indels 0; Gaps 0;  
 CC Qy 1 CUGGAUAGCUNGGAUCACCCUG 25  
 DB 33 CTGGATAGCTTGATCACCCCTG 9  
 RESULT 10  
 ID ADR13967/c  
 AC ADR13967 standard; cDNA; 198 BP.  
 XX  
 AC ADR13967;  
 XX DT 23-SEP-2004 (first entry)  
 XX DB Human NOGO-66 cDNA.  
 XX  
 KW ss; gene; human; myelin-associated glycoprotein; MAG; neural growth; neural regeneration; apoptosis; amyotrophic lateral sclerosis; Alzheimer's disease; Parkinson's disease; Huntington's disease; multiple sclerosis; Creutzfeldt-Jacob disease; kuru; multiple system atrophy; Lou Gehrig's disease; progressive supranuclear palsy.  
 XX OS Homo sapiens.  
 XX  
 PH Location/Qualifiers  
 1. .198  
 /tag= a  
 /partial  
 /product= "NOGO-66"  
 /notes "No start and stop codons given"  
 XX PN US2004121341-A1.  
 XX  
 PD 24-JUN-2004.  
 XX  
 PP 20-DEC-2002; 2002US-00327213.  
 XX  
 PR 20-DEC-2002; 2002US-00327213.  
 XX  
 PA (FILB/) FILBIN M.T.  
 PA (DOME/) DOMENICONI M.  
 PA (CAOZ/) CAO Z.  
 XX PT Filbin MT, Domeniconi M, Cao Z;  
 XX DR WPI; 2004-79666/45.  
 DR p-PSDB; ADR13968.  
 XX  
 PT New myelin-associated glycoprotein (MAG) derivative comprises a mutation in or flanking MAG Ig-like domain 5 (Ig5), excluding the MAG derivative (d1-3)-Fc, useful promoting neural growth and regeneration.  
 XX PS Disclosure; SEQ ID NO 10; 81pp; English.  
 CC The invention relates to a myelin-associated glycoprotein (MAG) derivative comprising a mutation in or flanking MAG Ig-like domain 5 (Ig5), excluding the MAG derivative (d1-3)-Fc, where the mutation reduces or eliminates the ability of the derivative to regulate neurite outgrowth as compared to endogenous or soluble MAG without eliminating binding to neuronal surfaces. The inhibitors of MAG are useful for promoting neural growth and regeneration. They are also useful for treating neural degeneration associated with injuries, disorders, or diseases. The disorder, disease, or condition is associated with apoptosis or results from a demyelinating disease and includes amyotrophic lateral sclerosis, Alzheimer's disease, Parkinson's disease, Huntington's disease, multiple sclerosis, Creutzfeldt-Jacob disease, kuru, multiple system atrophy, amyotrophic lateral sclerosis (Lou Gehrig's disease), or progressive supranuclear palsy. The present sequence represents the human NOGO-66 cDNA.  
 CC Sequence 198 BP; 56 A; 36 C; 49 G; 57 T; 0 U; 0 Other;  
 CC Query Match 100.0%; Score 25; DB 12; Length 198;  
 CC Best Local Similarity 72.0%; Pred. No. 0.079; Mismatches 0; Indels 0; Gaps 0;  
 CC Matches 18; Conservative 7; Mismatches 0; Indels 0; Gaps 0;  
 CC Qy 1 CUGGAUAGCUNGGAUCACCCUG 25  
 DB 33 CTGGATAGCTTGATCACCCCTG 9  
 RESULT 11  
 ID AAV23697/c  
 AC AAV23697 standard; cDNA; 261 BP.  
 XX  
 AC AAV23697;  
 XX DT 24-JUL-1998 (first entry)  
 XX  
 KW Human NSP1P protein coding sequence fragment.  
 KW NSP1P; neuroendocrine-specific protein-like protein; human; gene therapy; neurodegenerative disease; amyotrophic lateral sclerosis; cancer; ss.  
 XX OS Homo sapiens.  
 XX PN WO9806841-A2.  
 XX PD 19-FEB-1998.  
 XX PR 24-JUL-1997; 97WO-US013469.  
 XX PR 12-AUG-1997; 96US-00700607.  
 XX PA (INCYT-) INCYTE PHARM INC.  
 XX

PI Bandman O, Au-Young J, Goli SK, Hillman J;  
 XX  
 DR WPI; 1998-159533/14.  
 XX Human neuro-endocrine-specific protein-like protein - useful for  
 PT diagnosis, monitoring and treatment of cancer and neuro-degenerative  
 PT disease.  
 XX  
 PS Disclosure; Page 45; 73pp; English.  
 XX This sequence encodes a human neuroendocrine-specific protein-like  
 CC protein (NSPLP) of the invention. Recombinant cells transformed with the  
 DNA are used to express the NSPLP proteins, which are used to treat  
 cancer and neurodegenerative diseases such as amyotrophic lateral  
 sclerosis. Also antisense nucleic acids and antagonists of NSPLP can be  
 used to inhibit activity of the NSPLP proteins. Antibodies specific for  
 NSPLP are used for diagnosis and monitoring treatment of diseases  
 CC associated with NSPLP expression, in usual immunoassays, and to isolate  
 NSPLP from natural sources. The NSPLP proteins, or their fragments can  
 also be used in drug screening to identify NSPLP antagonists. The nucleic  
 acid can be used diagnostically and for monitoring treatment (in  
 hybridisation or amplification assays), to isolate closely related  
 sequences; in gene therapy for both sense and antisense applications  
 CC (including use of ribozymes) and for mapping the natural genomic sequence.  
 XX Sequence 261 BP; 62 A; 59 C; 56 G; 67 T; 0 U; 17 Other;  
 CC 100.0%; Score 25; DB 2; Length 261;  
 XX Best Local Similarity 72.0%; Pred. No. 0.082; 0; Mismatches  
 Matches 18; Conservative 18; MisMatches 0; Indels 0; Gaps 0;  
 OY 1 CUGGAUAGCUGGGAUCACCCUUG 25  
 DB 124 CTGGATAGCTGGATCACCCCTG 100

RESULT 12  
 AAX41193/C  
 ID AAX41193 standard; cDNA; 404 BP.  
 XX  
 AC AAX41193;  
 XX DT 17-JUN-1999 (first entry)  
 XX DB Human secreted protein 5' EST SEQ ID NO:137.  
 XX  
 KW Human; Secreted protein; EST; expressed sequence tag; diagnosis;  
 KW forensic; gene therapy; chromosome mapping; signal peptide;  
 KW upstream regulatory sequence; cytokine activity; cell proliferation;  
 KW differentiation; haematoopoiesis regulation; tissue growth regulation;  
 KW reproductive hormone regulation; chemotactic; chemokinetic; haemostatic;  
 KW thrombolytic; anti-inflammatory; tumour inhibition; ds.  
 OS Homo sapiens.  
 XX  
 PN WO9906548-A2.  
 XX  
 PD 11-FEB-1999.  
 XX 31-JUL-1998; 9BWO-IB001222.  
 XX  
 PR 01-AUG-1997; 97US-00905135.  
 XX  
 PA (GBST ) GENSET.  
 XX Dumas Milne Edwards J, Ducleart A, Lacroix B;  
 PI  
 XX WPI; 1999-153778/13.  
 DR P-PSDB; MAY12360.  
 XX  
 PT New nucleic acids encoding human secreted proteins - obtained from cDNA  
 PT libraries prepared from e.g. liver, ovary, brain, prostate, kidney, lung,  
 PT umbilical cord, placenta and colon tissue.

XX  
 PS Claim 1; Page 319; 824pp; English.  
 XX  
 AAX41094 to AAX41347 represent 5' expressed sequence tags (ESTs) for  
 CC human secreted proteins, and encode the proteins given in AAY1261 to  
 CC AAY154, respectively. The proteins given represent the signal peptide  
 CC and an N-terminal fragment of a secreted protein. The nucleic acid  
 CC sequences can be used for producing secreted human gene products. They  
 CC can also be used to develop products for diagnosis and therapy. The  
 CC proteins obtained may have cytokine activity, cell  
 CC proliferation/differentiation activity, haematopoiesis regulating  
 CC activity, tissue growth regulating activity, haemostatic and  
 CC regulating activity, chemotactic/ chemokinetic activity, anti-inflammatory  
 CC and thrombolytic activity, receptor/ ligand activity, anti-inflammatory  
 CC activity, tumour inhibition activity or other activities. The products  
 CC can be used in forensic, gene therapy and chromosome mapping procedures.  
 CC The sequences can also be used for obtaining corresponding promoter  
 CC sequences. The nucleic acids encoding the signal peptide can be used for  
 CC directing extracellular secretion of a polypeptide or the insertion of a  
 CC polypeptide into a membrane, or importing a polypeptide into a cell  
 XX Sequence 404 BP; 110 A; 75 C; 108 G; 111 T; 0 U; 0 Other;  
 CC  
 OY 1 CUGGAUAGCUGGGAUCACCCUUG 25  
 DB 347 CTGGATAGCTGGATCACCCCTG 323  
 OY  
 RESULT 13  
 AAF90323/C  
 ID AAF90323 standard; cDNA; 600 BP.  
 XX  
 AC AAF90323;  
 XX DT 23-JUL-2001 (first entry)  
 XX DB Human NOGO-C cDNA.  
 XX  
 KW Nogo-C; human; chromosome 2p21; neuropathy; spinal injury; brain injury;  
 KW stroke; neuronal degeneration; Alzheimer's disease; Parkinson's disease;  
 KW neurodegenerative disorder; psychiatric disorder; developmental disorder;  
 KW neuroprotective; nootropic; neuroleptic; antiparkinsonian;  
 KW cerebroprotective; neuroleptic; diagnosis; therapy; ss.  
 OS Homo sapiens.  
 XX  
 PN WO200136631-A1.  
 XX  
 PD 25-MAY-2001.  
 XX 14-NOV-2000; 2000WO-GB004345.  
 XX 15-NOV-1999; 99GB-00024995.  
 PR 24-JAN-2000; 2000GB-00001550.  
 XX  
 PA (SMIK ) SMITHKLINE BECHAM PLC.  
 XX  
 PI Michalovich D, Prinjha R;  
 XX  
 DR WPI; 2001-343822/36.  
 PR P-PSDB; AAB8234B.  
 XX  
 PT New polypeptide designated NOGO-C is a splice variant of the human NOGO  
 PT gene and may be useful in the treatment of neural disorders including  
 PT Alzheimer's and Parkinson's diseases.  
 XX  
 PS Claim 1; Page 25; 25pp; English.  
 XX  
 CC The present sequence is that of cDNA encoding human NOGO-C (see

CC AAB82348). NOGO-C is a novel splice variant of the human NOGO gene on  
 CC chromosome 22p1. 2 Other splice variant, NOGO-A and NOGO-B, have  
 CC previously been identified. The invention provides NOGO-C polypeptides  
 CC and polynucleotides, and methods for producing such polypeptides by  
 CC recombinant techniques. Also disclosed are methods for utilising NOGO-C  
 CC polypeptides and polynucleotides in the treatment of diseases including  
 CC neuropathies, spinal injury, brain injury, stroke, neuronal degeneration,  
 CC for example Alzheimer's disease and Parkinson's disease, neuromuscular  
 CC disorders, psychiatric disorders and developmental disorders. Also  
 CC provided are methods for identifying agonists and antagonists for use in  
 CC treating conditions associated with NOGO-C imbalance, and diagnostic  
 CC assays for detecting diseases associated with inappropriate NOGO-C  
 CC activity or levels  
 XX Sequence 600 BP; 161 A; 113 C; 144 G; 182 T; 0 U; 0 Other;  
 SQ Query Match 100.0%; Score 25; DB 4; Length 600;  
 Best Local Similarity 72.0%; Pred. No. 0.093; Mismatches 0; Indels 0; Gaps 0;  
 Matches 18; Conservative 7; Mismatches 0; Indels 0; Gaps 0;  
 Qy 1 CUGGAAUAGGUAGGAUCACCCUG 25  
 Db 216 CTTGGATAGCTTGATCACCCCTTG 192

RESULT 14  
 ABN96987/c  
 ID ABL89601 standard; DNA; 639 BP.  
 XX AC ABL89601;  
 AC ABL89601/  
 AC ABL89601/c  
 XX DT 24-MAY-2002 (first entry)  
 DB Human polynucleotide SEQ ID NO 163.  
 XX KW Cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral;  
 KW antiallergic; hepatotropic; antidiabetic; antiinflammatory; antilulcer;  
 KW antiviral; anticonvulsant; antibacterial; antifungal; antiarasitic;  
 KW cardiot; gene therapy; cancer; immune disorder; cardiovascular disorder;  
 KW neurological disease; infection; human; secreted protein; gene; ss.  
 XX OS Homo sapiens.  
 XX PN WO200190304-A2.  
 XX PD 29-NOV-2001.  
 XX PR 18-MAY-2001; 2001WO-US016450.  
 XX PR 19-MAY-2000; 2000US-0205515P.  
 XX PR (HUMA-) HUMAN GENOME SCI INC.  
 XX PT Birse CB, Rosen CA;  
 XX PT WPI; 2002-122018/16.  
 DR P2PSDB; ABB89192.  
 XX PS Novel 1405 isolated polypeptides, useful for diagnosis, treatment and  
 PT prevention of neural, immune system, muscular, reproductive,  
 PT gastrointestinal, pulmonary, cardiovascular, renal and proliferative  
 PT disorders.  
 XX PS Claim 4; SEQ ID NO 163; 2081pp + Sequence Listing; English.  
 XX The invention relates to novel genes (HLB89449-AB90853) and proteins  
 CC (ABB8040-ABB90444) useful for preventing, treating or ameliorating  
 CC medical conditions e.g. by protein or gene therapy. The genes are  
 CC isolated from a range of human tissues disclosed in the specification.  
 CC The nucleic acids, proteins, antibodies and (antagonists are useful in  
 CC the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and  
 CC ovarian cancer and other cancers of the adrenal gland, bone, bone marrow,  
 CC breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune  
 CC disorders e.g. Addison's disease, allergies, autoimmune haemolytic  
 CC anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease,  
 CC multiple sclerosis, rheumatoid arthritis and ulcerative colitis; (c)  
 CC cardiovascular disorders such as myocardial ischaemic; (d) wound healing  
 CC (e) neurological diseases e.g. cerebral anoxia and epilepsy; and (f)  
 CC infection diseases such as viral, bacterial, fungal and parasitic  
 CC infections. Note: The sequence data for this patent did not form part of  
 CC the printed specification, but was obtained in electronic format directly  
 CC from WIPO at [fp.wipo.int/pub/published\\_pct\\_sequences](http://fp.wipo.int/pub/published_pct_sequences)

CC for this patent did not form part of the printed specification, but was  
 CC obtained in electronic format directly from WIPO at  
 CC [fp.wipo.int/pub/published\\_pct\\_sequences](http://fp.wipo.int/pub/published_pct_sequences)  
 XX SQ Sequence 639 BP; 138 A; 114 C; 149 G; 147 T; 0 U; 91 Other;  
 CC Query Match 100.0%; Score 25; DB 6; Length 639;  
 Best Local Similarity 72.0%; Pred. No. 0.094; Mismatches 0; Indels 0; Gaps 0;  
 Matches 18; Conservative 7; Mismatches 0; Indels 0; Gaps 0;  
 Qy 1 CUGGAAUAGGUAGGAUCACCCUG 25  
 Db 415 CTTGGATAGCTTGATCACCCCTTG 391